## SEQUENCE LISTING

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<110> Bayer Aktiengesellschaft
<120> Ultraspiracle (USP) protein from Heliothis virescens
<130> Le A 34 771
<140>
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<150> DE 100 36 469.1
<151> 2000-07-25
<160> 2
<170> PatentIn Ver. 2.1
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                                                                   96
Ile Asn Trp Ala Arg Pro Leu Pro Pro Gly Gln Gln Gln Pro Met
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                                                      30
                                 25
acg cct acg tcg ccc gga aac atg ctt caa ccg atg gct acg ccg tct
Thr Pro Thr Ser Pro Gly Asn Met Leu Gln Pro Met Ala Thr Pro Ser
         35
                             40
                                                  45
aac tta ccg act gtc gac tgc tca ctc gat att caa tgg cta aac ttg
Asn Leu Pro Thr Val Asp Cys Ser Leu Asp Ile Gln Trp Leu Asn Leu
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                         55
gag gga ggt ttt atg tcg ccg atg tca ccg ccg gag atg aag cca gac
Glu Gly Gly Phe Met Ser Pro Met Ser Pro Pro Glu Met Lys Pro Asp
 65
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acg gcg Thr Ala									288
ttc aag Phe Lys						_			336
tgt tct Cys Ser							_		384
agt tgt Ser Cys 130								_	432
tta acg Leu Thr 145									480
cag agg Gln Arg								_	528
ggc atg Gly Met									576
ggt acg Gly Thr						_			624
tca atc Ser Ile 210									672
gaa gag Glu Glu 225						-	_		720
aag ttc Lys Phe									768
ata gcg Ile Ala								_	816

gag Glu										864
ctg Leu 290										912
cga Arg										960
ctt Leu										1008
cag Gln									_	1056
ctg Leu										1104
aag Lys 370										1152
caa Gln						-	_	_	_	1200
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atg atg

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Thr Pro Thr Ser Pro Gly Asn Met Leu Gln Pro Met Ala Thr Pro Ser

35 40 45

Asn Leu Pro Thr Val Asp Cys Ser Leu Asp Ile Gln Trp Leu Asn Leu 50 55 60

Glu Gly Gly Phe Met Ser Pro Met Ser Pro Pro Glu Met Lys Pro Asp
65 70 75 80

Thr Ala Met Leu Asp Gly Leu Arg Asp Asp Ser Thr Pro Pro Ala 85 90 95

Phe Lys Asn Tyr Pro Pro Asn His Pro Leu Ser Gly Ser Lys His Leu 100 105 110

Cys Ser Ile Cys Gly Asp Arg Ala Ser Gly Lys His Tyr Gly Val Tyr 115 120 125

Ser Cys Glu Gly Cys Lys Gly Phe Phe Lys Arg Thr Val Arg Lys Asp 130 135 140

Leu Thr Tyr Ala Cys Arg Glu Glu Arg Asn Cys Ile Ile Asp Lys Arg 145 150 155 160

Gln Arg Asn Arg Cys Gln Tyr Cys Arg Tyr Gln Lys Cys Leu Ala Cys 165 170 175

Gly Met Lys Arg Glu Ala Val Gln Glu Glu Arg Gln Arg Ala Ala Arg 180 185 190

Gly Thr Glu Asp Ala His Pro Ser Ser Ser Val Gln Val Gln Glu Leu

195 200 205

Ser Ile Glu Arg Leu Leu Glu Met Glu Ser Leu Val Ala Asp Pro Ser 210 215 220

Glu Glu Phe Gln Phe Leu Arg Val Gly Pro Asp Ser Asn Val Pro Pro 225 230 235 240

Lys Phe Arg Ala Pro Val Ser Ser Leu Cys Gln Ile Gly Asn Lys Gln
245 250 255

Ile Ala Ala Leu Val Val Trp Ala Arg Asp Ile Pro His Phe Ser Gln 260 265 270

Leu Glu Met Glu Asp Gln Ile Leu Leu Ile Lys Gly Ser Trp Asn Glu 275 280 285

Leu Leu Phe Ala Ile Ala Trp Arg Ser Met Glu Phe Leu Thr Glu 290 295 300

Glu Arg Asp Gly Val Asp Gly Thr Gly Asn Arg Thr Thr Ser Pro Pro 305 310 315 320

Gln Leu Met Cys Leu Met Pro Gly Met Thr Leu His Arg Asn Ser Ala 325 330 335

Leu Gln Ala Gly Val Gly Gln Ile Phe Asp Arg Val Leu Ser Glu Leu 340 345 350

Ser Leu Lys Met Arg Thr Leu Arg Val Asp Gln Ala Glu Tyr Val Ala 355 360 365

Leu Lys Ala Ile Ile Leu Leu Asn Pro Asp Val Lys Gly Leu Lys Asn 370 375 380

Arg Gln Glu Val Glu Val Leu Arg Glu Lys Met Phe Leu Cys Leu Asp 385 390 395 400

Glu Tyr Cys Arg Arg Ser Arg Ser Ser Glu Glu Gly Arg Phe Ala Ala 405 410 415

Leu Leu Leu Arg Leu Pro Ala Leu Arg Ser Ile Ser Leu Lys Ser Phe 420 425 430

Glu His Leu Phe Phe Phe His Leu Val Ala Asp Thr Ser Ile Ala Gly
435 440 445

Tyr Ile Arg Asp Ala Leu Arg Asn His Ala Pro Pro Ile Asp Thr Asn

Met Met